

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/8/4,634	<u>/</u>
Source:	IFNO-	
Date Processed bwSTIC:	1118/04	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221 Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to (EFFECTIVE 12/0 N03):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

	10 1010/C=>(C
ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER!
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8'Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
Invalid <213> Response	Per 1.823 of Sequence Rules, the only <u>valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species) <220> <223> section is required when <213> response is Unknown or is Artificial Sequence</u>
Usc of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
3 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/814,634

DATE: 04/08/2004 TIME: 10:04:11

Input Set : A:\SEQLIST FINAL.TXT

Output Set: N:\CRF4\04082004\J814634.raw

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4 <110> APPLICANT: CHENEVAL, Dominique
              KASTELIC, Tania
              Novation Pharmaceuticals Inc.
      8 <120> TITLE OF INVENTION: Assay for identifying Compounds Which
             Affect Stability of mRNA
     12 <130> FILE REFERENCE: 793-104CIP
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/814,634
C--> 14 <141> CURRENT FILING DATE: 2004-04-01
     14 <150> PRIOR APPLICATION NUMBER: 09/869,159
                                                               Does Not Comply
     15 <151> PRIOR FILING DATE: 1999-12-23
                                                               Corrected Diskette Needed
     17 <160> NUMBER OF SEQ ID NOS: 30
     19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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     22 <211> LENGTH: 1105
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Homo Sapiens
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     28 tgtccattta tagaataatg tgggaagaaa caaacccgtt ttatgattta ctcattatcg 120
     29 cettttgaca getgtgetgt aacacaagta gatgeetgaa ettgaattaa tecacacate 180
     30 agtaatgtat tetatetete tttacatttt ggtetetata etacattatt aatgggtttt 240
     31 gtgtactgta aagaatttag ctgtatcaaa ctagtgcatg aatagattct ctcctgatta 300
     32 tttatcacat agccccttag ccagttgtat attattcttg tggtttgtga cccaattaag 360
     33 tectaettta catatgettt aagaategat gggggatget teatgtgaac gtgggagtte 420
     34 agetgettet ettgeetaag tatteettte etgateacta tgeattttaa agttaaacat 480
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     36 agattgctgc ttctgctata tttgtgatat aggaattaag aggatacaca cgtttgtttc 600.
     37 ttcgtgcctg ttttatgtgc acacattagg cattgagact tcaagetttt ctttttttgt 660
     38 ccacgtatet ttgggtettt gataaagaaa agaateeetg tteattgtaa geaettttae 720
     39 gggggggtg gggaggggtg ctctgctggt cttcaattac caagaattct ccaaaacaat 780
     40 tttctgcagg atgattgtac agaatcattg cttatgacat gatcgctttc tacactgtat 840
     41 tacataaata aattaaataa aataaccccg ggcaagactt ttctttgaag gatgactaca 900
     42 gacattaaat aatcgaagta attttgggtg gggagaagag gcagattcaa ttttctttaa 960
     43 ccagtctgaa gtttcattta tgatacaaaa gaagatgaaa atggaagtgg caatataagg 1020
     44 ggatgaggaa ggcatgcctg gacaaaccct tcttttaaga tgtgtcttca atttgtataa 1080
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54 tagaaataat atgcattgtc agtgatgtac catgaaacaa agctgcaggc tgtttaagaa 120

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/814,634

DATE: 04/08/2004 TIME: 10:04:11

Input Set : A:\SEQLIST FINAL.TXT

Output Set: N:\CRF4\04082004\J814634.raw

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57 tttacattat taagaaaaaa agatttattt atttaagaca gtcccatcaa aactcctgtc 300
58 tttggaaatc cgaccactaa ttgccaagca ccgcttcgtg tggctccacc tggatgttct 360
59 gtgcctgtaa acatagattc gctttccatg ttgttggccg gatcaccatc tgaagagcag 420
60 acggatggaa aaaggacctg atcattgggg aagctggctt tctggctgct ggaggctggg 480
61 gagaaggtgt tcattcactt gcatttettt geeetggggg etgtgatatt aacagaggga 540
62 qqqttcctqt qqqqqaaqt ccatqcctcc ctqqcctqaa qaagaqactc tttqcatatq 600
63 actcacatga tgcatacctg gtgggaggaa aagagttggg aacttcagat ggacctagta 660
64 cccactgaga tttccacgcc gaaggacagc gatgggaaaa atgcccttaa atcataggaa 720
65 agtatttttt taagctacca attgtgccga gaaaagcatt ttagcaattt atacaatatc 780
66 atccagtace ttaagecetg attgtgtata tteatatatt ttggataege accececaae 840
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78 aaaataacac acatataaac atcacacaca cagacagaca cacacacaca caacaattaa 180
79 cagtetteag geaaaaegte gaateageta tttaetgeea aagggaaata teatttattt 240
80 tttacattat taagaaaaaa agatttattt atttaagaca gtcccatcaa aactcctgtc 300
81 tttggaaatc cgaccactaa ttgccaaqca ccqcttcgtq tggctccacc tggatgttct 360
82 gtgcctgtaa acatagatte getttecatg ttgttggccg gateaceate tgaagageag 420
83 acggatggaa aaaggacctg atcattgggg aagctggctt tctggctgct ggaggctggg 480
84 gagaaggtgt teatteactt geatttettt geectggggg etgtgatatt aacagaggga 540
85 gggtteetgt ggggggaagt ceatgeetee etggeetgaa gaagagaete tttgeatatg 600
86 acteacatga tgeatacetg gtgggaggaa aagagttggg aactteagat ggacetagta 660
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90 <211> LENGTH: 688
91 <212> TYPE: DNA
92 <213> ORGANISM: Homo Sapiens
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97 gaggagcaaa agctcatttc tgaagaggac ttgttgcgga aacgacgaga acagttgaaa 180
98 cacaaacttg aacagctacg gaactcttgt gcgtaaggaa aagtaaggaa aacgattcct 240
99 totgacagaa atgtootgag caatcaccta tgaacttgtt toaaatgcat gatcaaatgc 300
100 aacctcacaa ccttggctga gtcttgagac tgaaagattt agccataatg taaactgcct 360
101 caaattggac tttgggcata aaagaacttt tttatgctta ccatcttttt tttttcttta 420
102 acagatttgt atttaagaat tgtttttaaa aaattttaag atttacacaa tgtttctctg 480
103 taaatattgc cattaaatgt aaataacttt aataaaacgt ttatagcagt tacacagaat 540
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108 <210> SEQ ID NO: 5
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RAW SEQUENCE LISTING DATE: 04/08/2004 PATENT APPLICATION: US/10/814,634 TIME: 10:04:11

Input Set : A:\SEQLIST FINAL.TXT

Output Set: N:\CRF4\04082004\J814634.raw

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/814,634
Input Set: A:\SEQLIST FINAL.TXT

Output Set: N:\CRF4\04082004\J814634.raw

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164 cttatcttga acagatattt aattttgeta acactcaget etgeceteee egateeeetg 720
165 getecceage acacatteet ttgaaataag gttteaatat acatetacat actatatata 780
166 tatatttggc aacttgtatt tgtgtgtata tatatatata tatgtttatg tatatatgtg 840
167 attotqataa aatagacatt gotattotqt tttttatatq taaaaacaaa acaagaaaaa 900
168 atagagaatt ctacatacta aatctctctc cttttttaat tttaatattt gttatcattt 960
169 atttattggt gctactgttt atccgtaata attgtgggga aaagatatta acatcacgtc 1020
170 tttgtctcta gtgcagtttt tcgagatatt ccgtagtaca tatttatttt taaacaacga 1080
171 caaagaaata cagaacatat q
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174 <211> LENGTH: 168
175 <212> TYPE: DNA
176 <213> ORGANISM: Homo Sapiens
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180 tatatatata aaaataaata tetetattit atatatata aatatatata tiettititti 120
181 aaattaacag tgctaatgtt attggtgtct tcactggatg aacatatg
                                                            Invalide (2137 Response)

Response (2137 Response)

Response has to be eithernown

aagttgg has tificial Univenown

Trivalide Artificial Speciesso

Response OR Genus | Speciesso

Response
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184 <211> LENGTH: 33
185 <212> TYPE: DNA
186 <213> ORGANISM: Primer
188 <400> SEQUENCE: 9
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192 <211> LENGTH: 30
193 <212> TYPE: DNA _
194 <213> ORGANISM: Primer
196 <400> SEQUENCE: 10
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                                                                                                               lease see item #10
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201 <212> TYPE: DNA
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216 <211> LENGTH: 28
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218 <213> ORGANISM: Primer
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/814,634

DATE: 04/08/2004 TIME: 10:04:11

Same errors Input Set : A:\SEQLIST FINAL.TXT Output Set: N:\CRF4\04082004\J814634.raw 225 <212> TYPE: DNA 226 <213> ORGANISM: Primer 228 <400> SEQUENCE: 14 28 229 ccatatggct caatgatata tttgccag 231 <210> SEQ ID NO: 15 232 <211> LENGTH: 32 233 <212> TYPE: DNA 234 <213> ORGANISM: Primer 236 <400> SEQUENCE: 15 237 ageggeeget eggagetttt ttgeeetgeg tg 32 239 <210> SEQ ID NO: 16 240 <211> LENGTH: 28 241 <212> TYPE: DNA 242 <213> ORGANISM: Primer 244 <400> SEQUENCE: 16 245 ccatatgaag caaactttat ttctcgcc 28 247 <210> SEQ ID NO: 17 248 <211> LENGTH: 31 249 <212> TYPE: DNA 250 <213> ORGANISM: Primer 252 <400> SEQUENCE: 17 253 agcggccgct gaggaggacg aacatccaac c 31 255 <210> SEQ ID NO: 18 256 <211> LENGTH: 27 257 <212> TYPE: DNA 258 <213> ORGANISM Primer 260 <400> SEQUENCE: 18 27 261 ccatatggtg aagtttattt cagaacc 263 <210> SEQ ID NO: 19 264 <211> LENGTH: 30 265 <212> TYPE: DNA 266 <213> ORGANISM: Primer 268 <400> SEQUENCE: 19 30 269 agcggccgct aaagagagct gtacccagag 271 <210> SEQ ID NO: 20 272 <211> LENGTH: 32 273 <212> TYPE: DNA 274 <213> ORGANISM: Primer 276 <400> SEQUENCE: 20 277 aacatatgtt ctgtatttct ttgtcgttgt tt 32 279 <210> SEQ ID NO: 21 280 <211> LENGTH: 32 281 <212> TYPE: DNA 282 <213> ORGANISM: (Primer 284 <400> SEQUENCE: 21 32 285 tgcggccgca ttgctgtgct ttggggattc cc 287 <210> SEQ ID NO: 22 288 <211> LENGTH: 33 289 <212> TYPE: DNA The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/814,634

DATE: 04/08/2004 TIME: 10:04:12

Input Set : A:\SEQLIST FINAL.TXT
Output Set: N:\CRF4\04082004\J814634.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date